

21	24	2.4	1320	3	US-08-461-	Sequence 8, Application	3.15e-02
22	24	2.4	1620	3	US-08-611-	Sequence 10, Application	3.15e-02
23	24	2.4	2167	3	US-08-611-	Sequence 9, Application	3.15e-02
24	24	2.4	2668	3	US-08-611-	Sequence 11, Application	3.15e-02
25	23	2.3	66	4	PCT-US55-1	Sequence 93, Application	1.19e-01
26	23	2.3	75	4	PCT-US55-1	Sequence 99, Application	1.19e-01
27	23	2.3	81	4	PCT-US55-1	Sequence 92, Application	1.19e-01
28	22	2.2	65	1	US-08-471-	Sequence 145, Application	4.37e-01
29	22	2.2	66	1	PCT-US55-1	Sequence 93, Application	4.37e-01
30	22	2.2	68	1	US-08-471-	Sequence 143, Application	4.37e-01
31	22	2.2	69	1	US-08-471-	Sequence 142, Application	4.37e-01
32	22	2.2	74	4	PCT-US55-1	Sequence 94, Application	4.37e-01
33	22	2.2	105	1	US-07-665-	Sequence 13, Application	4.37e-01
34	22	2.2	2380	3	US-08-672-	Sequence 3, Application	4.37e-01
35	21	2.1	66	1	US-08-672-	Sequence 144, Application	1.56e-001
36	21	2.1	92	3	US-08-553-	Sequence 4, Application	1.56e-001
37	21	2.1	242	1	US-08-273-	Sequence 1, Application	1.56e-001
38	21	2.1	2277	3	US-08-676-	Sequence 2, Application	1.56e-001
39	21	2.1	2277	2	US-08-676-	Sequence 2, Application	1.56e-001
40	20	2.0	59	4	PCT-US55-1	Sequence 95, Application	5.36e-001
41	20	2.0	59	4	PCT-US55-1	Sequence 95, Application	5.36e-001
42	20	2.0	85	2	US-08-538-	Sequence 1, Application	5.36e-001
43	20	2.0	85	4	PCT-US44-0	Sequence 26, Application	5.36e-001
44	20	2.0	242	4	PCT-US44-0	Sequence 1, Application	5.36e-001
45	20	2.0	1106	4	PCT-US55-0	Sequence 3, Application	5.36e-001
ALIGNMENTS							
AC	AC	1	US-08-232-463-14	STANDARD	DNR: UNC: 7218 BP.		
DT	DT	1	US-08-232-463				
DE	Sequence 14, Application	US/08232463					
CC	Sequence 14, Application	US/08232463					
CC	Patent No. 5,670,367						
CC	GENERAL INFORMATION:						
CC	APPLICANT: DORNER, F.						
CC	APPLICANT: SCHEFFLINGER, F.						
CC	APPLICANT: FALKNER, F. G.						
CC	TITLE OF INVENTION: RECOMBINANT FOWLPOX VIRUS						
CC	NUMBER OF SEQUENCES: 52						
CC	CORRESPONDENCE ADDRESS:						
CC	ADDRESSEE: Foley & Lardner						
CC	STREET: 1800 Diagonal Road, Suite 500						
CC	CITY: Alexandria						
CC	STATE: VA						
CC	COUNTRY: USA						
CC	ZIP: 22313-0299						
CC	COMPUTER READABLE FORM:						
CC	MEDIUM TYPE: FLOPPY DISK						
CC	COMPUTER: IBM PC compatible						
CC	OPERATING SYSTEM: PC-DOS/MS-DOS						
CC	SOFTWARE: Patentin Release #1.0, Version #1.25						
CC	CURRENT APPLICATION DATA:						
CC	APPLICATION NUMBER: US/08/232,463						
CC	FILING DATE: 26-AUG-1991						
CC	FILING DATE:						
CC	CLASSIFICATION: 435						
CC	PRIOR APPLICATION DATA:						
CC	APPLICATION NUMBER: US/07/935,313						
CC	FILING DATE:						
CC	APPLICATION NUMBER: EP 91 114 300.6						
CC	FILING DATE: 26-AUG-1991						
CC	ATTORNEY/AGENT INFORMATION:						
CC	NAME: BENT, Stephen A.						
CC	REGISTRATION NUMBER: 29,768						
CC	REFERENCE/DOCKET NUMBER: 30472/114 IMMU						
CC	TELECOMMUNICATION INFORMATION:						
CC	TELEPHONE: (703)856-9300						
CC	TELEX: 899149						
CC	TELEX: (703)683-4109						
CC	INFORMATION FOR SEQ ID NO: 14:						
CC	SEQUENCE CHARACTERISTICS:						



CC INFORMATION FOR SEQ ID NO: 14:  
 CC SEQUENCE CHARACTERISTICS:  
 CC LENGTH: 7218 base Pairs  
 CC TYPE: nucleic acid  
 CC STRANDEDNESS: single  
 CC TOPOLOGY: linear  
 CC IMMEDIATE SOURCE:  
 CC CLONE: PTZOPT-F1S  
 CC SQ SEQUENCE 7218 BP; 1944 A; 1491 C; 1486 G; 1929 T; 368 OTHER.  
 CC Query Match 3.8%; Score 38; DB 3; Length 965;  
 CC Best Local Similarity 17.5%; Pred. No. 3 35e-11;  
 CC Matches 30; Conservative 74; Mismatches 66; Indels 1; Gaps 1;  
 CC RESULT 4  
 ID US-08-3888-672A-22 STANDARD; DNA; UNC; 965 BP.  
 AC XXXXXX  
 DT Sequence 22, Application US/08388672A  
 CC Sequence 22, Application US/08388672A  
 CC Patent No. 5795961  
 CC GENERAL INFORMATION:  
 CC APPLICANT: Wallace, T. Paul  
 CC APPLICANT: Harris, William J.  
 CC APPLICANT: Carr, Frank J.  
 CC APPLICANT: Old, Lloyd J.  
 CC APPLICANT: Welt, Sydney  
 CC APPLICANT: Kitamura, Kunio  
 CC TITLE OF INVENTION: Recombinant Human Anti-Lewis B  
 CC NUMBER OF SEQUENCES: 25  
 CC CORRESPONDENCE ADDRESS:  
 CC ADDRESSEE: Felfe and Lynch  
 CC STREET: 805 Third Avenue  
 CC CITY: New York  
 CC STATE: New York  
 CC COUNTRY: U.S.A.  
 CC ZIP: 10022  
 CC COMPUTER READABLE FORM:  
 CC MEDIUM TYPE: FLOPPY disk  
 CC COMPUTER: IBM PC compatible  
 CC OPERATING SYSTEM: PC-DOS/MS-DOS  
 CC SOFTWARE: PatentIn Release #1.0, version #1.30  
 CC CURRENT APPLICATION DATA:  
 CC APPLICATION NUMBER: US/08/238-163  
 CC FILING DATE: 03-MAY-1994  
 CC CLASSIFICATION: 800  
 CC ATTORNEY/AGENT INFORMATION:  
 CC NAME: Bastian, Kevin L.  
 CC REGISTRATION NUMBER: 34-774  
 CC REFERENCE/DOCKET NUMBER: 2307-E-540  
 CC TELECOMMUNICATION INFORMATION:  
 CC TELEPHONE: (415) 543-9600  
 CC TELEFAX: (415) 543-5043  
 CC INFORMATION FOR SEQ ID NO: 5:  
 CC SEQUENCE CHARACTERISTICS:  
 CC LENGTH: 215 base Pairs  
 CC TYPE: nucleic acid  
 CC STRANDEDNESS: single  
 CC TOPOLOGY: unknown  
 CC MOLECULE TYPE: protein  
 CC FEATURE:  
 CC NAME/KEY: misc feature  
 CC LOCATION: 1..215  
 CC OTHER INFORMATION: /standard\_name= "Deduced amino acid  
 CC SEQUENCE 215 BP; 15 A; 8 C; 25 G; 26 T; 141 OTHER.  
 CC Query Match 3.2%; Score 32; DB 1; Length 215;







DE Sequence 98, Application PC/TUS9511934  
 CC Sequence 98, Application PC/TUS9511934  
 CC GENERAL INFORMATION:  
 CC APPLICANT: Cytogen Corporation  
 CC TITLE OF INVENTION: Antigen Binding Peptides (Abtides) From  
 CC NUMBER OF SEQUENCES: 103  
 CC CORRESPONDENCE ADDRESS:  
 CC ADDRESSEE: Pennie & Edmonds  
 CC STREET: 115 Avenue of the Americas  
 CC CITY: New York  
 CC STATE: New York  
 CC COUNTRY: USA  
 CC ZIP: 10036  
 CC COMPUTER READABLE FORM:  
 CC MEDIUM TYPE: Floppy disk  
 CC COMPUTER: IBM PC compatible  
 CC OPERATING SYSTEM: PC-DOS/MS-DOS  
 CC SOFTWARE: PatentIn Release #1.0, Version #1.30  
 CC CURRENT APPLICATION DATA:  
 CC APPLICATION NUMBER: PCT/US95/11934  
 CC FILING DATE: 20-SEP-1995  
 CC CLASSIFICATION:  
 CC ATTORNEY/AGENT INFORMATION:  
 CC NAME: Misrock, S. Leslie  
 CC REFERENCE/DOCKET NUMBER: 1101-196-228  
 CC TELECOMMUNICATION INFORMATION:  
 CC TELEPHONE: 212-790-9090  
 CC TELEFAX: 212-669-8864/9741  
 CC TELEX: 66141 PENNIE  
 CC INFORMATION FOR SEQ ID NO: 145:  
 CC MOLECULE TYPE: DNA  
 CC LENGTH: 65 bases  
 CC TYPE: nucleic acid  
 CC STRANDEDNESS: single  
 CC TOPOLOGY: unknown  
 CC SEQUENCE 65 BP; 3 A; 3 C; 3 G; 2 T; 54 OTHER.  
 CC SQ  
 CC SEQUENCE TYPE: DNA (genomic)  
 CC SEQUENCE 81 BP; 6 A; 6 C; 4 G; 5 T; 60 OTHER.  
 CC  
 CC Query Match 2.5%; Score 25; DB 4; Length 81;  
 CC Best Local Similarity 10.4%; Pred. No. 8.13e-03;  
 CC Matches 7; Conservative 20; Mismatches 40; Indels 0; Gaps 0;  
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 CC Query Match 2.4%; Score 24; DB 1; Length 65;  
 CC Best Local Similarity 10.0%; Pred. No. 3.15e-02;  
 CC Matches 6; Conservative 18; Mismatches 36; Indels 0; Gaps 0;  
 CC  
 CC Sequence 143, Application US/08471052A  
 CC Sequence 143, Application US/08471052A  
 CC Patent No. 5625033  
 CC GENERAL INFORMATION:  
 CC APPLICANT: Kay, B. K.  
 CC APPLICANT: Fowlkes, D. M.  
 CC TITLE OF INVENTION: Totally Synthetic Affinity Reagents  
 CC NUMBER OF SEQUENCES: 166  
 CC CORRESPONDENCE ADDRESS:  
 CC ADDRESSEE: Pennie & Edmonds  
 CC STREET: 115 Avenue of the Americas  
 CC CITY: New York  
 CC STATE: New York  
 CC COUNTRY: U.S.A.  
 CC ZIP: 10036-2711  
 CC COMPUTER READABLE FORM:  
 CC MEDIUM TYPE: Floppy disk  
 CC COMPUTER: IBM PC compatible  
 CC OPERATING SYSTEM: PC-DOS/MS-DOS  
 CC SOFTWARE: PatentIn Release #1.0, Version #1.25  
 CC CURRENT APPLICATION DATA:  
 CC APPLICATION NUMBER: US/08471052A  
 CC FILING DATE: 06-JUNE-1995  
 CC CLASSIFICATION: 530  
 CC ATTORNEY/AGENT INFORMATION:  
 CC NAME: Misrock, S. Leslie  
 CC REFERENCE/DOCKET NUMBER: 18,872  
 CC TELECOMMUNICATION INFORMATION:  
 CC TELEPHONE: 212-790-9090  
 CC TELEFAX: 212-669-8864/9741  
 CC TELEX: 66141 PENNIE  
 CC INFORMATION FOR SEQ ID NO: 143:  
 CC SEQUENCE CHARACTERISTICS:  
 CC LENGTH: 68 bases

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 SQ      TYPE: nucleic acid  
 STRANDEDNESS: single  
 TOPOLOGY: unknown  
 MOLECULE TYPE: DNA  
 SEQUENCE 68 BP; 3 A; 3 C; 5

CC MOLECULE TYPE: DNA  
SQ SEQUENCE 68 BP; 3 A; 3 C; 5 G; 3 T; 54 OTHER.

TYPE: nucleic acid  
STRANDEDNESS: single  
TOPOLOGY: unknown

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Query Match          2.48;  Score 24;  DB 1;  Length 68;
Best Local Similarity 10.0%;  Pred. No. 3.15e-02;
Matches 6;  Conservative
Matches 6;  Strict
Mismatches 35;  Redels 0;  Gaps 0;  Gaps 0;

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Search completed: Fri Jan 14 07:05:14 2000  
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